

6-26-02

GP 1652
Box 580

In the United States Patent and Trademark Office

RECEIVED

JUN 28 2002

TECH CENTER 1600/2900

In re Application of:
Alberto I. Roca

Serial No. 09/358,103

Filed: 07/21/1999

For: Mutants of MAW Motifs of
RecA Protein Homologs, Methods
of Making Them, and Their Uses



§
§
§
§
§
§
§
§
§
§

Group Art Unit: 1652

Examiner: Christian Fronda

Atty. Docket: ROCA-01

22/f
J.G.J
7/3/02

SUPPLEMENTAL AMENDMENT

Honorable Commissioner of Patents
and Trademarks
Washington, D.C. 20231

Dear Sir:

In response to the Office Communication dated 05/21/2002:

Applicant submits herewith a computer readable form copy of the sequence listing, a paper copy of this sequence listing, and the inventor's declaration that the computer readable form and paper copies of the listing are the same. Please amend the application by entering this sequence listing.

Date: 6/21/02

Respectfully submitted,

R. Perry McConnell
Registration No. 38,239
R. Perry McConnell, P.C.
9001 Forest Crossing, Suite F
The Woodlands, Texas 77381
Telephone: 281-296-9200
Facsimile: 281-296-9393

RECEIVED

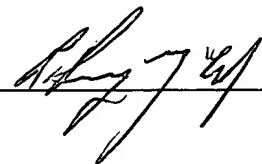
JUN 28 2002

CERTIFICATE OF MAILING VIA "EXPRESS MAIL" UNDER 37 C.F.R. § 1.10

TECH CENTER 1600/2900

I hereby certify that the above **Supplemental Amendment** is being mailed to the Assistant Commissioner of Patents, Washington, D.C. 20231, via the United States Postal Service Express Mail, on the 21st day of June, 2002.

Express Mail No.: EU099976831US





RECEIVED

JUN 28 2002

TECH CENTER 1600/2900

In the United States Patent and Trademark Office

In re Application of:
Alberto I. Roca

Serial No. 09/358,103

Filed: 07/21/1999

For: Mutants of MAW Motifs of
RecA Protein Homologs, Methods
of Making Them, and Their Uses



Group Art Unit: 1652

Examiner: R.A. Wax

Atty. Docket: ROCA-01

Declaration of Alberto I. Roca

I declare that the information recorded in computer readable form submitted with this amendment is identical to the written sequence listing. This amended listing contains no new matter over the content of the original application specification.

I declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

Date

1/13/00


Alberto I. Roca

SEQUENCE LISTING



RECEIVED

JUN 28 2002

TECH CENTER 1600/2900

<110> Roca, Alberto I

<120> Mutants of MAW Motifs of RecA Protein Homologs, Methods
of Making Them, and Their Uses

<130> RecA Homolog Protein & Mutants

<140> 09/358,103

<141> 1999-07-21

<150> 60/094,071

<151> 1998-07-24

<160> 3

<170> PatentIn Ver. 2.0

<210> 1

<211> 26

<212> PRT

<213> Escherichia coli

<220>

<221> NON_TER

<222> (1)

<220>

<221> NON_TER

<222> (26)..)

<220>

<221> HELIX

<222> (6)..(12)

<223> Alpha-helix B

<220>

<221> STRAND

<222> (22)..(26)

<223> Beta-strand 1

<220>

<221> SIMILAR

<222> (1)..(26)

<223> This structure is highly conserved across
bacterial RecA and homologous eukaryotic,
archaeal, and viral proteins; sequence below is

from E. coli RecA positions 40-65

<400> 1

Ile Ser Thr Gly Ser Leu Ser Leu Asp Ile Ala Leu Gly Ala Gly Gly
1 5 10 15

Leu Pro Met Gly Arg Ile Val Glu Ile Tyr
20 25

<210> 2

<211> 26

<212> PRT

<213> Escherichia coli

<220>

<221> NON_TER

<222> (1)

<220>

<221> NON_TER

<222> (26)

<220>

<221> HELIX

<222> (6)..(12)

<223> Alpha-helix B

<220>

<221> STRAND

<222> (22)..(26)

<223> Beta-strand 1

<220>

<221> SIMILAR

<222> (1)..(26)

<223> Non 'Xaa' residues are the invariant MAW-motif
residues in RecA and its homologs

<220>

<221> SITE

<222> (1)

<223> This site is not invariant across RecA homologs
and has observed to contain Ile, Phe, Met, Thr,
Val, Tyr, or Leu

<220>

<221> SITE

3'
Cent

<222> (2)
<223> This site is not invariant across RecA homologs
and has been observed to contain Ser, Gly, Lys,
Pro, Thr, or Arg

<220>
<221> SITE
<222> (3)
<223> This site is not invariant across RecA homologs
and has been observed to contain Thr, Ser, or Trp

<220>
<221> SITE
<222> (4)
<223> This site is not invariant across RecA homologs
and has been observed to contain Gly, Arg, Ala,
Lys, Asn, or Gln

<220>
<221> SITE
<222> (5)
<223> This site is not invariant across RecA homologs
and has been observed to contain Ser, Ala, Cys,
Ile, Asn, Asp, or Phe

<220>
<221> SITE
<222> (6)
<223> This site is not invariant across RecA homologs
and has been observed to contain Leu, Ile, Thr,
Val, Tyr, Lys, Gln, Asp, Pro, or Arg

<220>
<221> SITE
<222> (7)
<223> This site is not invariant across RecA homologs
and has been observed to contain Ser, Ala, Asp,
Gly, Leu, Met, Thr, Val, Tyr, Glu, Asn, or Gln

<220>
<221> SITE
<222> (8)
<223> This site is not invariant across RecA homologs
and has been observed to contain Leu, Ile, Val,
Phe, or Met

<220>
<221> SITE

3!
Cont

<222> (9)

<223> This site is not invariant across RecA homologs
and has been observed to contain Asp or Asn

<220>

<221> SITE

<222> (10)

<223> This site is not invariant across RecA homologs
and has been observed to contain Ile, Ala, Glu,
Gly, Leu, Asn, Gln, Arg, Ser, Thr, Val, Lys, or
Asp

<220>

<221> SITE

<222> (11)

<223> This site is not invariant across RecA homologs
and has been observed to contain Ala, Ile, Leu, or
Val

<220>

<221> SITE

<222> (12)

<223> This site is not invariant across RecA homologs
and has been observed to contain Leu, Met, or Thr

<220>

<221> SITE

<222> (13)

<223> This site is not invariant across RecA homologs
and has been observed to contain Gly, Gln, Ala,
Asn, or Ser

<220>

<221> SITE

<222> (14)

<223> This site is not invariant across RecA homologs
and has been observed to contain Ala, Ile, Ser,
Thr, Val, Gly, or Leu, and this site is not
present in some homologs

<220>

<221> SITE

<222> (17)

<223> This site is not invariant across RecA homologs
and has been observed to contain Leu, Phe, Ile,
Val, Tyr, or Met

<220>

31
Cont.

<221> SITE
<222> (18)
<223> This site is not invariant across RecA homologs
and has been observed to contain Pro, Glu, Met,
Phe, Gln, Arg, or Val

<220>
<221> SITE
<222> (19)
<223> This site is not invariant across RecA homologs
and has been observed to contain Met, Gly, Lys,
Arg, Thr, Ala, Pro, or Ser

<220>
<221> SITE
<222> (20)
<223> This site is not invariant across RecA homologs
and has been observed to contain Gly, Leu, Met,
Ala, His, Gln, or Arg

<220>
<221> SITE
<222> (21)
<223> This site is not invariant across RecA homologs
and has been observed to contain Arg, Gln, Ser,
Gly, Thr, or Val

<220>
<221> SITE
<222> (22)
<223> This site is not invariant across RecA homologs
and has been observed to contain Ile, Val, Ala,
Leu, or Met

<220>
<221> SITE
<222> (23)
<223> This site is not invariant across RecA homologs
and has been observed to contain Val, Ile, Thr, or
Tyr

<220>
<221> SITE
<222> (25)
<223> This site is not invariant across RecA homologs
and has been observed to contain Ile, Val, Ala,
Leu, Met, or Phe

gl
cont

<220>

<221> SITE

<222> (26)

<223> This site is not invariant across RecA homologs
and has been observed to contain Tyr, Phe, Ala,
Gly, or Val

<400> 2

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Gly
1 5 10 15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu Xaa Xaa
20 25

<210> 3

<211> 26

<212> PRT

<213> Escherichia coli

<220>

<221> NON_TER

<222> (1)

<220>

<221> NON_TER

<222> (26)

<220>

<221> HELIX

<222> (6)..(12)

<223> Alpha-helix B

<220>

<221> STRAND

<222> (22)..(26)

<223> Beta-strand 1

<220>

<221> SIMILAR

<222> (1)..(26)

<223> Non "Xaa" residues are the invariant and
semiconservative elements of the MAW motif in RecA
and its homologs

<220>

<221> SITE

<222> (2)

gl
Ant.

<223> This site is neither invariant nor
semiconservative across RecA homologs and has been
observed to contain Ser, Gly, Lys, Pro, Thr, or
Arg

<220>

<221> SITE

<222> (5)

<223> This site is neither invariant nor
semiconservative across RecA homologs and has been
observed to contain Ser, Ala, Cys, Ile, Asn, Asp,
or Phe

<220>

<221> SITE

<222> (6)

<223> This site is neither invariant nor
semiconservative across RecA homologs and has been
observed to contain Leu, Ile, Thr, Val, Tyr, Lys,
Gln, Asp, Pro, or Arg

<220>

<221> SITE

<222> (7)

<223> This site is neither invariant nor
semiconservative across RecA homologs and has been
observed to contain Ser, Ala, Asp, Gly, Leu, Met,
Thr, Val, Tyr, Glu, Asn, or Gln

<220>

<221> SITE

<222> (10)

<223> This site is neither invariant nor
semiconservative across RecA homologs and has been
observed to contain Ile, Ala, Glu, Gly, Leu, Asn,
Gln, Arg, Ser, Thr, Val, Lys, or Asp

<220>

<221> SITE

<222> (13)

<223> This site is neither invariant nor
semiconservative across RecA homologs and has been
observed to contain Gly, Gln, Ala, Asn, or Ser

<220>

<221> SITE

<222> (14)

<223> This site is neither invariant nor

31
Ant

semiconservative across RecA homologs and has been observed to contain Ala, Ile, Ser, Thr, Val, Gly, or Leu, and this site is not present in some homologs

<220>

<221> SITE

<222> (18)

<223> This site is neither invariant nor semiconservative across RecA homologs and has been observed to contain Pro, Glu, Met, Phe, Gln, Arg, or Val

<220>

<221> SITE

<222> (19)

<223> This site is neither invariant nor semiconservative across RecA homologs and has been observed to contain Met, Gly, Lys, Arg, Thr, Ala, Pro, or Ser

<220>

<221> SITE

<222> (21)

<223> This site is neither invariant nor semiconservative across RecA homologs and has been observed to contain Arg, Gln, Ser, Gly, Thr, or Val

<400> 3

Ile Xaa Thr Gly Xaa Xaa Xaa Leu Asp Xaa Ala Leu Xaa Xaa Gly Gly
1 5 10 15

Leu Xaa Xaa Gly Xaa Ile Val Glu Ile Tyr
20 25

